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From: Graser, Jennifer **AU 1645**
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011 N26 **8/21/24/05**

Ngo, J.T. et al., "Computational Complexity, Protein Structure Prediction, and the Levinthal Paradox",
The Protein Folding Problem and Tertiary Structure Prediction, 1994, Merz et al (eds), Birkhauser, Boston,
MA, pp. 433 and 492-495.

Mikayama et al, Molecular Cloning and Functional Expression of a cDNA encoding glycosylation-inhibiting
factor, November 1993, Proc. Nat. Acad. Sci., USA, vol 90: 10056-10060.
"human glycosylation-inhibiting factor (GIF) protein differs from human macrophage migration inhibitory factor
(MIF) by a single amino acid residue (Figure 1) and GIF is unable to carry out the function of MIF and MIF does not
demonstrate GIF bioactivity (ABST)."